

RAW SEQUENCE LISTING

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Application Serial Number: 10/582,061
Source: IFWP
Date Processed by STIC: 06/19/2006

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IFWP

RAW SEQUENCE LISTING

DATE: 06/19/2006

PATENT APPLICATION: US/10/582,061

TIME: 11:40:47

Input Set : A:\Herr-Band5.00955-03.txt

Output Set: N:\CRF4\06192006\J582061.raw

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3 <110> APPLICANT: University of Virginia Patent Foundation
4     Herr, John C
5     Sleight, Susan C.
7 <120> TITLE OF INVENTION: Band 5: A human testis specific protein
9 <130> FILE REFERENCE: 00955-03
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/582,061
C--> 11 <141> CURRENT FILING DATE: 2006-06-08
11 <150> PRIOR APPLICATION NUMBER: US 60/527,875
12 <151> PRIOR FILING DATE: 2003-12-08
14 <150> PRIOR APPLICATION NUMBER: PCT/US04/41440
15 <151> PRIOR FILING DATE: 2004-12-08
17 <160> NUMBER OF SEQ ID NOS: 9
19 <170> SOFTWARE: PatentIn version 3.3
21 <210> SEQ ID NO: 1
22 <211> LENGTH: 10
23 <212> TYPE: PRT
24 <213> ORGANISM: Mus musculus
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33 <211> LENGTH: 9
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44 <211> LENGTH: 13
45 <212> TYPE: PRT
46 <213> ORGANISM: Mus musculus
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57 <213> ORGANISM: Mus musculus
59 <400> SEQUENCE: 4
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65 <210> SEQ ID NO: 5
66 <211> LENGTH: 14

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68 <213> ORGANISM: Mus musculus
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84 caaggaaaac cttactctta gtcgtgagaa agtattttta aaagtcctca gaacctggac      120
86 tgagctgggt gaagcattag cgcttgaggg cgcggaattg gaaagcggct tttaaaggg      180
88 gttgtttaag cgccctcccc cctgcccctc gtttgggggt agaggatcct ctggcggtgc      240
90 tggcaagctg tgcgttctgg ccttcattgg gccgcatttt acactcttgc tggcagctct      300
92 tgccaactgc ctgtgtccag ggaggccctg catcaaagt gaccagtttg tgacagatgc      360
94 gctaaagact ttcgaaaaca cttacctgaa tgaccacctg ccacacgaca ttcacaaaaa      420
96 tgtaatgagg atggtgaacc atgaagtatc gagcttcggc gtagtcactt cggctgagga      480
98 ttctattttg ggggcccgtg acgagaacac actggaacaa gcaacctgga gttttctgaa      540
100 ggatctgaag cgtattacag acagtgactt aaaaggagag ctctttataa aggaactatt      600
102 gtggatgctt cgtcatcaaa aggacatctt taacaatctt gctagacagt tccaaaagga      660
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106 cgaaaagcag ttgcacattt gtcggaaatc cctagatttg ggagagcgcc acatagaggt      780
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114 ctgtgtgcta gataccatca accaaggtca tgccaccgtc atccgctaag atgtcacagt      1020
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118 cgagactcct gtccacgtga ctccacagac accaccgggg caggagccag agtcggagct      1140
120 gtaccgggag ctcgaccag agttgtacct ggagtcata cctacagtcg ctcaaatcc      1200
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126 aagtgcagaa gtcaaaccce ccgcgtcagg atccaagtc gatcagagtt tgtcccaaca      1380
128 gatgggattg aagaaggcct ctcaggcaga ttttaactct gactactctg gagataagag      1440
130 cgaggcaaca gaaaactaat aaagatttgt atttgactc      1479
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134 <211> LENGTH: 397
135 <212> TYPE: PRT
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144 Cys Pro Gly Arg Pro Cys Ile Lys Cys Asp Gln Phe Val Thr Asp Ala
145          20          25          30
148 Leu Lys Thr Phe Glu Asn Thr Tyr Leu Asn Asp His Leu Pro His Asp
149          35          40          45
152 Ile His Lys Asn Val Met Arg Met Val Asn His Glu Val Ser Ser Phe
153          50          55          60
156 Gly Val Val Thr Ser Ala Glu Asp Ser Tyr Leu Gly Ala Val Asp Glu

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157 65          70          75          80
160 Asn Thr Leu Glu Gln Ala Thr Trp Ser Phe Leu Lys Asp Leu Lys Arg
161          85          90          95
164 Ile Thr Asp Ser Asp Leu Lys Gly Glu Leu Phe Ile Lys Glu Leu Leu
165          100          105          110
168 Trp Met Leu Arg His Gln Lys Asp Ile Phe Asn Asn Leu Ala Arg Gln
169          115          120          125
172 Phe Gln Lys Glu Val Leu Cys Pro Asn Lys Cys Gly Val Met Ser Gln
173          130          135          140
176 Thr Leu Ile Trp Cys Leu Lys Cys Glu Lys Gln Leu His Ile Cys Arg
177 145          150          155          160
180 Lys Ser Leu Asp Cys Gly Glu Arg His Ile Glu Val His Arg Ser Glu
181          165          170          175
184 Asp Leu Val Leu Asp Cys Leu Leu Ser Trp His Arg Ala Ser Lys Gly
185          180          185          190
188 Leu Thr Asp Tyr Ser Phe Tyr Arg Val Trp Glu Asn Ser Ser Glu Thr
189          195          200          205
192 Leu Ile Ala Lys Gly Lys Glu Pro Tyr Leu Thr Lys Ser Met Val Gly
193          210          215          220
196 Pro Glu Asp Ala Gly Asn Tyr Arg Cys Val Leu Asp Thr Ile Asn Gln
197 225          230          235          240
200 Gly His Ala Thr Val Ile Arg Tyr Asp Val Thr Val Leu Pro Pro Lys
201          245          250          255
204 His Ser Glu Glu Asn Gln Pro Pro Asn Ile Ile Thr Gln Glu Glu His
205          260          265          270
208 Glu Thr Pro Val His Val Thr Pro Gln Thr Pro Pro Gly Gln Glu Pro
209          275          280          285
212 Glu Ser Glu Leu Tyr Pro Glu Leu His Pro Glu Leu Tyr Pro Glu Leu
213          290          295          300
216 Ile Pro Thr Val Ala Gln Asn Pro Glu Lys Lys Met Lys Thr Arg Leu
217 305          310          315          320
220 Leu Ile Leu Leu Thr Leu Gly Phe Val Val Leu Val Ala Ser Ile Ile
221          325          330          335
224 Ile Ser Val Leu His Phe Arg Lys Val Ser Ala Lys Leu Lys Asn Ala
225          340          345          350
228 Ser Asp Glu Val Lys Pro Thr Ala Ser Gly Ser Lys Ser Asp Gln Ser
229          355          360          365
232 Leu Ser Gln Gln Met Gly Leu Lys Lys Ala Ser Gln Ala Asp Phe Asn
233          370          375          380
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237 385          390          395
240 <210> SEQ ID NO: 8
241 <211> LENGTH: 350
242 <212> TYPE: PRT
243 <213> ORGANISM: Homo sapiens
245 <400> SEQUENCE: 8
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248 1          5          10          15
251 Leu Pro Ala Glu Gly Cys Val Ile Cys Asp Pro Ser Val Val Leu Ala

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252          20          25          30
255 Leu Lys Ser Leu Glu Lys Asp Tyr Leu Pro Gly His Leu Asp Ala Lys
256          35          40          45
259 His His Lys Ala Met Met Glu Arg Val Glu Asn Ala Val Lys Asp Phe
260          50          55          60
263 Gln Glu Leu Ser Leu Asn Glu Asp Ala Tyr Met Gly Val Val Asp Glu
264 65          70          75          80
267 Ala Thr Leu Gln Lys Gly Ser Trp Ser Leu Leu Lys Asp Leu Lys Arg
268          85          90          95
271 Ile Thr Asp Ser Asp Val Lys Gly Asp Leu Phe Val Lys Glu Leu Phe
272          100         105         110
275 Trp Met Leu His Leu Gln Lys Glu Thr Phe Ala Thr Tyr Val Ala Arg
276          115         120         125
279 Phe Gln Lys Glu Ala Tyr Cys Pro Asn Lys Cys Gly Val Met Leu Gln
280          130         135         140
283 Thr Leu Ile Trp Cys Lys Asn Cys Lys Lys Glu Val His Ala Cys Arg
284 145         150         155         160
287 Lys Ser Tyr Asp Cys Gly Glu Arg Asn Val Glu Val Pro Gln Met Glu
288          165         170         175
291 Asp Met Ile Leu Asp Cys Glu Leu Asn Trp His Gln Ala Ser Glu Gly
292          180         185         190
295 Leu Thr Asp Tyr Ser Phe Tyr Arg Val Trp Gly Asn Asn Thr Glu Thr
296          195         200         205
299 Leu Val Ser Lys Gly Lys Glu Ala Thr Leu Thr Lys Pro Met Val Gly
300          210         215         220
303 Pro Glu Asp Ala Gly Ser Tyr Arg Cys Glu Leu Gly Ser Val Asn Ser
304 225         230         235         240
307 Ser Pro Ala Thr Ile Asn Phe His Val Thr Val Leu Pro Lys Met
308          245         250         255
311 Ile Lys Glu Glu Lys Pro Ser Pro Asn Ile Val Thr Pro Gly Glu Ala
312          260         265         270
315 Thr Thr Glu Ser Ser Ile Ser Leu Gln Pro Leu Gln Pro Glu Lys Met
316          275         280         285
319 Leu Ala Ser Arg Leu Leu Gly Leu Leu Ile Cys Gly Ser Leu Ala Leu
320          290         295         300
323 Ile Thr Gly Leu Thr Phe Ala Ile Phe Arg Arg Arg Lys Val Ile Asp
324 305         310         315         320
327 Phe Ile Lys Ser Ser Leu Phe Gly Leu Gly Ser Gly Val Ala Glu Gln
328          325         330         335
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332          340         345         350
335 <210> SEQ ID NO: 9
336 <211> LENGTH: 1695
337 <212> TYPE: DNA
338 <213> ORGANISM: Homo sapiens
340 <400> SEQUENCE: 9
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343 agttccttgt ggagagggga tcgtccgtga ctcagacccc gtcgaaaagc tctcgatttc      120
345 acgcaaccaa ccttaccctc gtccccaaca caatccttga ttttaccctt gggaaaatta      180

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347	gcgtaggtgg	tcatgacccc	ggcttctctct	ttaaaagccc	tcagaacctg	gaccgggctc	240
349	ctcaatgaac	cctagtgtta	gaggccacaa	atcggaatc	aaggtcgggg	agcggctgtt	300
351	taggggtgcat	cactgttcag	gatgcacgg	ggctcctttt	tctaagcgga	agtgtgcgcc	360
353	ctgcaaattg	gaagcccgtt	agttttgggg	aagggatgaac	aagggccccc	taggtctaga	420
355	ggaccgcatg	gagtctcgct	taagtgttag	tggggtcctc	agttctagaa	atctgcgttc	480
357	tcctgctctc	ggtttggggg	gagagcacc	ttcctaacgg	tcttcgggaa	ctgtgcgcgc	540
359	cggtgcaat	ggggccgcat	tttaccctcc	tgtgtgcggc	gctggccggg	tgttgccttc	600
361	ctgccgaggg	gtgtgttata	tgtgaccggt	ctgtcgtgct	ggcgctaaag	tccttgagga	660
363	aagattacct	gcctggccac	ctggatgcca	agcatcaca	agccatgatg	gaaagggtag	720
365	agaatgccgt	gaaggatttc	caggaactgt	cgcttaatga	ggatgcctat	atgggggtcg	780
367	ttgatgaggc	cacactgcaa	aaggggtcct	ggagtttgct	gaaggatctg	aaacgcacat	840
369	cagacagtga	tgtaaaaggc	gatctctttg	tgaaggagct	atgttggtat	ttgcacttgc	900
371	aaaaggaaac	ctttgccacc	tatgttgctc	gattccaaaa	agaggcttat	tgtcccaaca	960
373	aatgtggtgt	gatgttgcaa	actctgatct	gggtgcaaga	ctgcaaaaag	gaggttcacg	1020
375	cttgtcgaaa	gtcctacgat	tgcggggagc	ggaatgtgga	agttcctcaa	atggaagaca	1080
377	tgatcctgga	ctgtgagtta	aactggcatc	aggcttcgga	aggcctcact	gattacagct	1140
379	tttacagggt	ttgggggaac	aatacggaga	ccttgggtgc	caaggggaaa	gaggccaccc	1200
381	tgaccaagcc	catggtgggt	ccagaggatg	caggcagcta	ccgctgcgag	ctgggctctg	1260
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385	aggaggaaaa	accttctcca	aatatcgtaa	ccccggggga	ggcgaccacg	gagtcgtcca	1380
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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/582,061

DATE: 06/19/2006

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L:11 M:270 C: Current Application Number differs, Replaced Current Application No

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date